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RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/390,207DATE: 09/21/1999  
TIME: 16:08:59

Input Set: I390207.RAW

This Raw Listing contains the General Information  
Section and up to first 5 pages.

P5  
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1 <110> APPLICANT: Thomason, Arlen  
2 Liu, Benxian  
3 <120> TITLE OF INVENTION: Fibroblast Growth Factor-Like Polypeptides  
4 <130> FILE REFERENCE: 99-371  
5 <140> CURRENT APPLICATION NUMBER: US/09/390,207  
6 <141> CURRENT FILING DATE: 1999-09-07  
7 <160> NUMBER OF SEQ ID NOS: 41  
8 <170> SOFTWARE: PatentIn Ver. 2.0  
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19 acctgaggac ccgagccatt g atg gac tgc gac gag acc ggg ttc gag cac 171  
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20 1 5 10  
21 tca gga ctg tgg gtt tct gtg ctg gct ctt ctg ctg gga gcc tgc 219  
22 Ser Gly Leu Trp Val Ser Val Leu Ala Gly Leu Leu Gly Ala Cys  
23 15 20 25  
24 cag gca cac ccc atc cct gac tcc agt cct ctc ctg caa ttc ggg ggc 267  
25 Gln Ala His Pro Ile Pro Asp Ser Ser Pro Leu Leu Gln Phe Gly Gly  
26 30 35 40  
27 caa gtc cgg cag cgg tac ctc tac aca gat gat gcc cag cag aca gaa 315  
28 Gln Val Arg Gln Arg Tyr Leu Tyr Thr Asp Asp Ala Gln Gln Thr Glu  
29 45 50 55  
30 gcc cac ctg gag atc agg gag gat ggg acg gtg ggg ggc gct gct gac 363  
31 Ala His Leu Glu Ile Arg Glu Asp Gly Thr Val Gly Gly Ala Ala Asp  
32 60 65 70  
33 cag agc ccc gaa agt ctc ctg cag aaa gcc ttg aag ccg gga gtt 411  
34 Gln Ser Pro Glu Ser Leu Leu Gln Leu Lys Ala Leu Lys Pro Gly Val  
35 75 80 85 90  
36 att caa atc ttg gga gtc aag aca tcc agg ttc ctg tgc cag cgg cca 459  
37 Ile Gln Ile Leu Gly Val Lys Thr Ser Arg Phe Leu Cys Gln Arg Pro  
38 95 100 105  
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40 Asp Gly Ala Leu Tyr Gly Ser Leu His Phe Asp Pro Glu Ala Cys Ser  
41 110 115 120  
42 ttc cgg gag ctg ctt ctt gag gac gga tac aat gtt tac cag tcc gaa 555  
43 Phe Arg Glu Leu Leu Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu  
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 59   Arg Ser Pro Ser Tyr Ala Ser  
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 62   agagtccag aggaggataa gaatgagcat gtgtgagtgt ctgagggaa acatggcagc   921  
 63   tgtttgtct cccttggccc ggacaatccc ctctacaccc cccctcacgt gttccgaggg   981  
 64   tcctggcttc ccactgggcc tcactttttt cttttctttt cttttctttt ttttgagacg   1041  
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 81   Glu Asp Gly Thr Val Gly Gly Ala Ala Asp Gln Ser Pro Glu Ser Leu  
 82                       65                       70                       75                       80  
 83   Leu Gln Leu Lys Ala Leu Lys Pro Gly Val Ile Gln Ile Leu Gly Val  
 84                       85                       90                       95  
 85   Lys Thr Ser Arg Phe Leu Cys Gln Arg Pro Asp Gly Ala Leu Tyr Gly  
 86                       100                       105                       110  
 87   Ser Leu His Phe Asp Pro Glu Ala Cys Ser Phe Arg Glu Leu Leu Leu  
 88                       115                       120                       125  
 89   Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu Ala His Gly Leu Pro Leu  
 90                       130                       135                       140  
 91   His Leu Pro Gly Asn Lys Ser Pro His Arg Asp Pro Ala Pro Arg Gly  
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 93   Pro Ala Arg Phe Leu Pro Leu Pro Gly Leu Pro Pro Ala Pro Pro Glu  
 94                       165                       170                       175

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110 1 5 10 15  
111 ctg ctg ctg gct gtc ttc ctg ctg ggg gtc tac caa gca tac ccc atc 96  
112 Leu Leu Leu Ala Val Phe Leu Leu Gly Val Tyr Gln Ala Tyr Pro Ile  
113 20 25 30  
114 cct gac tcc agc ccc ctc ctc cag ttt ggg ggt caa gtc cgg cag agg 144  
115 Pro Asp Ser Ser Pro Leu Leu Gln Phe Gly Gly Gln Val Arg Gln Arg  
116 35 40 45  
117 tac ctc tac aca gat gac gac caa gac act gaa gcc cac ctg gag atc 192  
118 Tyr Leu Tyr Thr Asp Asp Asp Gln Asp Thr Glu Ala His Leu Glu Ile  
119 50 55 60  
120 agg gag gat gga aca gtg gta ggc gca gca cac cgc agt cca gaa agt 240  
121 Arg Glu Asp Gly Thr Val Val Gly Ala Ala His Arg Ser Pro Glu Ser  
122 65 70 75 80  
123 ctc ctg gag ctc aaa gcc ttg aag cca ggg gtc att caa atc ctg ggt 288  
124 Leu Leu Glu Leu Lys Ala Leu Lys Pro Gly Val Ile Gln Ile Leu Gly  
125 85 90 95  
126 gtc aaa gcc tct agg ttt ctt tgc caa cag cca gat gga gct ctc tat 336  
127 Val Lys Ala Ser Arg Phe Leu Cys Gln Gln Pro Asp Gly Ala Leu Tyr  
128 100 105 110  
129 gga tcg cct cac ttt gat cct gag gcc tgc agc ttc aga gaa ctg ctg 384  
130 Gly Ser Pro His Phe Asp Pro Glu Ala Cys Ser Phe Arg Glu Leu Leu  
131 115 120 125  
132 ctg gag gac ggt tac aat gtg tac cag tct gaa gcc cat ggc ctg ccc 432  
133 Leu Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu Ala His Gly Leu Pro  
134 130 135 140  
135 ctg cgt ctg cct cag aag gac tcc cca aac cag gat gca aca tcc tgg 480  
136 Leu Arg Leu Pro Gln Lys Asp Ser Pro Asn Gln Asp Ala Thr Ser Trp  
137 145 150 155 160  
138 gga cct gtg cgc ttc ctg ccc atg cca ggc ctg ctc cac gag ccc caa 528  
139 Gly Pro Val Arg Phe Leu Pro Met Pro Gly Leu Leu His Glu Pro Gln  
140 165 170 175  
141 gac caa gca gga ttc ctg ccc cca gag ccc cca gat gtg ggc tcc tct 576  
142 Asp Gln Ala Gly Phe Leu Pro Pro Glu Pro Pro Asp Val Gly Ser Ser  
143 180 185 190  
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Input Set: I390207.RAW

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157 20 25 30  
158 Pro Asp Ser Ser Pro Leu Leu Gln Phe Gly Gly Gln Val Arg Gln Arg  
159 35 40 45  
160 Tyr Leu Tyr Thr Asp Asp Asp Gln Asp Thr Glu Ala His Leu Glu Ile  
161 50 55 60  
162 Arg Glu Asp Gly Thr Val Val Gly Ala Ala His Arg Ser Pro Glu Ser  
163 65 70 75 80  
164 Leu Leu Glu Leu Lys Ala Leu Lys Pro Gly Val Ile Gln Ile Leu Gly  
165 85 90 95  
166 Val Lys Ala Ser Arg Phe Leu Cys Gln Gln Pro Asp Gly Ala Leu Tyr  
167 100 105 110  
168 Gly Ser Pro His Phe Asp Pro Glu Ala Cys Ser Phe Arg Glu Leu Leu  
169 115 120 125  
170 Leu Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu Ala His Gly Leu Pro  
171 130 135 140  
172 Leu Arg Leu Pro Gln Lys Asp Ser Pro Asn Gln Asp Ala Thr Ser Trp  
173 145 150 155 160  
174 Gly Pro Val Arg Phe Leu Pro Met Pro Gly Leu Leu His Glu Pro Gln  
175 165 170 175  
176 Asp Gln Ala Gly Phe Leu Pro Pro Glu Pro Pro Asp Val Gly Ser Ser  
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190 20 25 30  
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192 35 40 45  
193 Pro Glu Ser Leu Leu Gln Leu Lys Ala Leu Lys Pro Gly Val Ile Gln  
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RAW SEQUENCE LISTING  
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Input Set: I390207.RAW

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199           85          90          95
200 Glu Leu Leu Leu Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu Ala His
201           100         105         110
202 Gly Leu Pro Leu His Leu Pro Gly Asn Lys Ser Pro His Arg Asp Pro
203           115         120         125
204 Ala Pro Arg Gly Pro Ala Arg Phe Leu Pro Leu Pro Gly Leu Pro Pro
205           130         135         140
206 Ala Pro Pro Glu Pro Pro Gly Ile Leu Ala Pro Gln Pro Pro Asp Val
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209           165         170         175
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220           20          25          30
221 Leu Glu Ile Arg Glu Asp Gly Thr Val Val Gly Ala Ala His Arg Ser
222           35          40          45
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224           50          55          60
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226           65          70          75          80
227 Ala Leu Tyr Gly Ser Pro His Phe Asp Pro Glu Ala Cys Ser Phe Arg
228           85          90          95
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231 Gly Leu Pro Leu Arg Leu Pro Gln Lys Asp Ser Pro Asn Gln Asp Ala
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233 Thr Ser Trp Gly Pro Val Arg Phe Leu Pro Met Pro Gly Leu Leu His
234           130         135         140
235 Glu Pro Gln Asp Gln Ala Gly Phe Leu Pro Pro Glu Pro Pro Asp Val
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244 <213> ORGANISM: Mus musculus

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Please Note:  
 Use of n and/or Xaa have been detected in the Sequence Listing. Please review the  
 Sequence Listing to ensure that a corresponding explanation is presented in the <220> to  
 <223> fields of each sequence which presents at least one n or Xaa.

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VERIFICATION SUMMARY  
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Line ? Error/Warning

Original Text

60 W Invalid/Missing Amino Acid Numbering  
935 W "N" or "Xaa" used: Feature required

205

210

ggaaggaaaa aagcggccgc aacannnnnn nnn